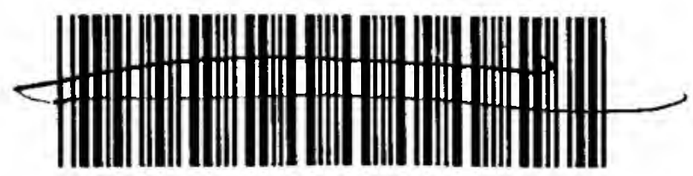
0420

#2



OIPE

RAW SEQUENCE LISTING DATE: 02/19/2002
PATENT APPLICATION: US/10/066,960 TIME: 12:40:15

Input Set : N:\Crf3\RULE60\10066960.txt
Output Set: N:\CRF3\02192002\J066960.raw

SEQUENCE LISTING

(1) GENERAL INFORMATION:
(i) APPLICANT: PARMA, et al.
(ii) TITLE OF INVENTION: HIGH AFFINITY NUCLEIC ACID LIGANDS TO LECTINS
(iii) NUMBER OF SEQUENCES: 390
(iv) CORRESPONDENCE ADDRESS:
(A) ADDRESSEE: Swanson & Bratschun, L.L.C.
(B) STREET: 8400 E. Prentice Avenue, Suite 200
(C) CITY: Englewood
(D) STATE: Colorado
(E) COUNTRY: USA
(F) ZIP: 80111
(v) COMPUTER READABLE FORM:
(A) MEDIUM TYPE: Diskette, 3 1/2 diskette, 1.44 MB
(B) COMPUTER: IBM pc compatible
(C) OPERATING SYSTEM: MS-DOS
(D) SOFTWARE: WordPerfect 6.0
(vi) CURRENT APPLICATION DATA:
C--> **(A) APPLICATION NUMBER: US/10/066,960**
C--> **(B) FILING DATE: 04-Feb-2002**
(C) CLASSIFICATION:
(vii) PRIOR APPLICATION DATA:
(A) APPLICATION NUMBER: 08/952,793
(B) FILING DATE: 1999-DEC-03
(A) APPLICATION NUMBER: PCT/US96/09455
(B) FILING DATE: 05-JUNE-1995
(A) APPLICATION NUMBER: 08/479,724
(B) FILING DATE: 07-JUNE-1995
(A) APPLICATION NUMBER: 08/472,256
(B) FILING DATE: 07-JUNE-1995
(A) APPLICATION NUMBER: 08/472,255
(B) FILING DATE: 07-JUNE-1995
(A) APPLICATION NUMBER: 08/477,829
(B) FILING DATE: 07-JUNE-1995
(viii) ATTORNEY/AGENT INFORMATION:
(A) NAME: Barry J. Swanson
(B) REGISTRATION NUMBER: 33,215
(C) REFERENCE/DOCKET NUMBER: NEX40C/PCT
(ix) TELECOMMUNICATION INFORMATION:
(A) TELEPHONE: (303) 793-3333
(B) TELEFAX: (303) 793-3433
(2) INFORMATION FOR SEQ ID NO: 1:

RAW SEQUENCE LISTING DATE: 02/19/2002
PATENT APPLICATION: US/10/066,960 TIME: 12:40:15

Input Set : N:\Crf3\RULE60\10066960.txt
Output Set: N:\CRF3\02192002\J066960.raw

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 98 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear
W--> **(ii) MOLECULE TYPE: RNA**
(ix) FEATURE:
(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine
(ix) FEATURE:
(D) OTHER INFORMATION: All U's are 2'-NH2 uracil
(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
GGGAAAAGCG AAUCAUACAC AAGANNNNNN NNNNNNNNNN NNNNNNNNNN 50
NNNNNNNNNN NNNNNNNNNN NNNNGCUCCG CCAGAGACCA ACCGAGAA 98
(2) INFORMATION FOR SEQ ID NO: 2:
(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 41 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear
W--> **(ii) MOLECULE TYPE: RNA**
(ix) FEATURE:
(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine
(ix) FEATURE:
(D) OTHER INFORMATION: All U's are 2'-NH2 uracil
(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
UAAUACGACU CACUAUAGGG AAAAGCGAAU CAUACACAAG A 41
(2) INFORMATION FOR SEQ ID NO: 3:
(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 24 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear
W--> **(ii) MOLECULE TYPE: RNA**
(ix) FEATURE:
(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine
(ix) FEATURE:
(D) OTHER INFORMATION: All U's are 2'-NH2 uracil
(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
UUCUCGGUUG GUCUCUGGCG GAGC 24
(2) INFORMATION FOR SEQ ID NO: 4:
(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 96 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear
W--> **(ii) MOLECULE TYPE: RNA**
(ix) FEATURE:
(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine
(ix) FEATURE:

RAW SEQUENCE LISTING DATE: 02/19/2002
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Input Set : N:\Crf3\RULE60\10066960.txt
Output Set: N:\CRF3\02192002\J066960.raw

(D) OTHER INFORMATION: All U's are 2'-NH2 uracil
(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
GGGAAAAGCG AAUCAUACAC AAGAAUGGUU GGCCUGGGCG CAGGCUUCGA 50
AGACUCGGCG GGAACGGGAA UGGCUCCGCC AGAGACCAAC CGAGAA 96

(2) INFORMATION FOR SEQ ID NO: 5:
(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 98 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear
W--> **(ii) MOLECULE TYPE: RNA**
(ix) FEATURE:
(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine
(ix) FEATURE:
(D) OTHER INFORMATION: All U's are 2'-NH2 uracil
(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:
GGGAAAAGCG AAUCAUACAC AAGACAGGCA CUGAAAACUC GGCGGGAACG 50
AAAGUAGUGC CGACUCAGAC GCGUGCUCCG CCAGAGACCA ACCGAGAA 98

(2) INFORMATION FOR SEQ ID NO: 6:
(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 91 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear
W--> **(ii) MOLECULE TYPE: RNA**
(ix) FEATURE:
(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine
(ix) FEATURE:
(D) OTHER INFORMATION: All U's are 2'-NH2 uracil
(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:
GGGAAAAGCG AAUCAUACAC AAGAAGUCUG GCCAAAGACU CGGCGGGAAC 50
GUAAAACGGC CAGAAUUGCU CCGCCAGAGA CCAACCGAGA A 91

(2) INFORMATION FOR SEQ ID NO: 7:
(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 94 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear
W--> **(ii) MOLECULE TYPE: RNA**
(ix) FEATURE:
(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine
(ix) FEATURE:
(D) OTHER INFORMATION: All U's are 2'-NH2 uracil
(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:
GGGAAAAGCG AAUCAUACAC AAGAGUAGGA GGUUCCAUCA CCAGGACUCG 50
GCGGGAACGG AAGGUGAUGS GCUCCGCCAG AGACCAACCG AGAA 94

(2) INFORMATION FOR SEQ ID NO: 8:
(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 95 base pairs

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Input Set : N:\Crf3\RULE60\10066960.txt
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(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear
W--> **(ii) MOLECULE TYPE: RNA**
(ix) FEATURE:
(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine
(ix) FEATURE:
(D) OTHER INFORMATION: All U's are 2'-NH2 uracil
(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:
GGGAAAAGCG AAUCAUACAC AAGAACAAGG AUCGAUGGCG AGCCGGGGAG 50
GGCUCGGCGG GAACGAAAUC UGCUCCGCCA GAGACCAACC GAGAA 95

(2) INFORMATION FOR SEQ ID NO: 9:
(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 97 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear
W--> **(ii) MOLECULE TYPE: RNA**
(ix) FEATURE:
(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine
(ix) FEATURE:
(D) OTHER INFORMATION: All U's are 2'-NH2 uracil
(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:
GGGAAAAGCG AAUCAUACAC AAGAUUGGGC AGGCAGAGCG AGACCGGGGG 50
CUCGGCGGGA ACGGAACAGG AAUGCUCCGC CAGAGACCAA CCGAGAA 97

(2) INFORMATION FOR SEQ ID NO: 10:
(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 97 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear
W--> **(ii) MOLECULE TYPE: RNA**
(ix) FEATURE:
(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine
(ix) FEATURE:
(D) OTHER INFORMATION: All U's are 2'-NH2 uracil
(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:
GGGAAAAGCG AAUCAUACAC AAGAAAGGGA UGGGAUUGGG ACGAGCGGCC 50
AAGACUCGGC GGGAACGAAG GGUGCUCCGC CAGAGACCAA CCGAGAA 97

(2) INFORMATION FOR SEQ ID NO: 11:
(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 96 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear
W--> **(ii) MOLECULE TYPE: RNA**
(ix) FEATURE:
(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine
(ix) FEATURE:

(D) OTHER INFORMATION: All U's are 2'-NH2 uracil

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:

GGGAAAAGCG AAUCAUACAC AAGACUCGGC GGGAACGAAA GUGUCAUGGU 50

AGCAAGUCCA AUGGUGGACU CUGCUCCGCC AGAGACCAAC CGAGAA 96

(2) INFORMATION FOR SEQ ID NO: 12:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 98 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

W--> **(ii) MOLECULE TYPE: RNA**

(ix) FEATURE:

(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine

(ix) FEATURE:

(D) OTHER INFORMATION: All U's are 2'-NH2 uracil

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 12:

GGGAAAAGCG AAUCAUACAC AAGACUCGGC GGGAACGUGA AGUGGGUAGG 50

UAGCUGAAGA CGGUCUGGGC GCCAGCUCCG CCAGAGACCA ACCGAGAA 98

(2) INFORMATION FOR SEQ ID NO: 13:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 99 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

W--> **(ii) MOLECULE TYPE: RNA**

(ix) FEATURE:

(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine

(ix) FEATURE:

(D) OTHER INFORMATION: All U's are 2'-NH2 uracil

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 13:

GGGAAAAGCG AAUCAUACAC AAGAAAGGGA UGGGAUUGGG ACGAGCGGCC 50

AAGACUCGGC GGGAACGAAG GGUCCGCUCC GCCAGAGACC AACCGAGAA 99

(2) INFORMATION FOR SEQ ID NO: 14:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 98 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

W--> **(ii) MOLECULE TYPE: RNA**

(ix) FEATURE:

(D) OTHER INFORMATION: All C's are 2'-NH2 cytosine

(ix) FEATURE:

(D) OTHER INFORMATION: All U's are 2'-NH2 uracil

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 14:

GGGAAAAGCG AAUCAUACAC AAGACUCGGC GGGAACGAAG UGUGUGAGUA 50

ACGAUCACUU GGUACUAAAA GCCCGCUCCG CCAGAGACCA ACCGAGAA 98

(2) INFORMATION FOR SEQ ID NO: 15:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 100 base pairs

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Input Set : N:\Crf3\RULE60\10066960.txt
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L:20 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]
L:21 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]
L:56 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:56 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=1
L:71 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:71 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=2
L:85 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:85 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=3
L:99 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:99 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=4
L:114 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:114 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=5
L:129 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:129 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=6
L:144 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:144 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=7
L:159 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:159 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=8
L:174 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:174 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=9
L:189 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:189 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=10
L:204 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:204 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=11
L:219 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:219 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=12
L:234 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:234 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=13
L:249 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:249 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=14
L:264 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:264 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=15
L:279 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:279 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=16
L:295 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:295 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=17
L:310 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:310 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=18
L:325 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:325 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=19
L:340 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:340 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=20
L:355 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:355 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=21
L:370 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:370 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=22
L:385 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:385 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=23

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Input Set : **N:\Crf3\RULE60\10066960.txt**
Output Set: **N:\CRF3\02192002\J066960.raw**

L:400 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:400 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=24
L:415 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:415 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=25
L:430 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:430 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=26
L:445 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:445 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=27
L:460 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:460 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=28
L:475 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:475 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=29
L:490 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:490 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=30
L:505 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:505 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=31
L:520 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:520 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=32
L:535 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:535 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=33
L:550 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:550 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=34
L:565 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:565 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=35
L:580 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:580 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=36
L:595 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:595 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=37
L:610 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:610 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=38
L:625 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:625 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=39
L:640 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:640 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=40
L:655 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:655 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=41
L:670 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:670 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=42
L:685 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:685 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=43
L:700 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:700 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=44
L:715 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:715 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=45
L:730 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:730 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=46
L:745 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]
L:745 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=47
L:760 M:220 C: Keyword misspelled or invalid format, [(ii) MOLECULE TYPE:]

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Input Set : N:\Crf3\RULE60\10066960.txt
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L:760 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=48
L:775 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=49
L:790 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=50